

Read mapping stats

Species	Read Length	Mapping	Library	Raw fragments	Complexity	Unique	Unique Splices	Multi	Multi Splices	Fraction mapped
hg20	2x75	STAR-2.5.3a GENCODE V25	AALE-KRAS-exoRNeasy-DKEL010C-7	2,800,128	0.61	3,845,133	727,209	409,979	96,189	0.91
hg20	2x75	STAR-2.5.3a GENCODE V25	AALE-KRAS-exotic-DKEL010F-13	2,446,934	0.79	3,077,588	174,976	584,610	51,340	0.79
hg20	2x75	STAR-2.5.3a GENCODE V25	AALE-PBABE-exoRNeasy-DKEL010D-8	2,956,271	0.61	4,065,022	910,560	348,358	118,414	0.92
hg20	2x75	STAR-2.5.3a GENCODE V25	AALE-PBABE-exotic-DKEL010E-12	446,268	0.93	676,037	35,303	34,169	4,873	0.84

#	Exonic:	Intergenic:	Intronic:
AALE-KRAS-exoRNeasy-DKEL010C-7	0.73	0.11	0.17
AALE-KRAS-exotic-DKEL010F-13	0.22	0.33	0.45
AALE-PBABE-exoRNeasy-DKEL010D-8	0.77	0.09	0.14
AALE-PBABE-exotic-DKEL010E-12	0.20	0.33	0.47

Coverage of genes; >1000bp

